

Introduction to Spatial Point Pattern Analysis

by

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References:

Ripley, B.D. (1981). *Spatial Statistics*. Wiley, New York.
Diggle, P.J. (2003). *Statistical Analysis of Spatial Point Patterns*, 2nd Ed. Oxford University Press, London.
Upton G.J.G., and Fingleton, B. (1985). *Spatial Data Analysis by Example*. Wiley, New York.
Waller, L.A., and Gotway, C.A. (2004). *Applied Spatial Statistics for Public Health Data*. Wiley, New York.
Møller, J., and Waagepetersen, R.P. (2004). *Statistical Inference and Simulation for Spatial Point Processes*. Chapman and Hall/CRC, Boca Raton, FL.

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Definition: A *spatial point pattern* is comprised of the locations of events.

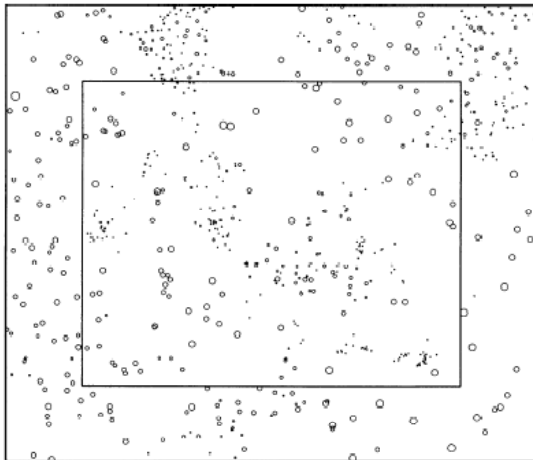
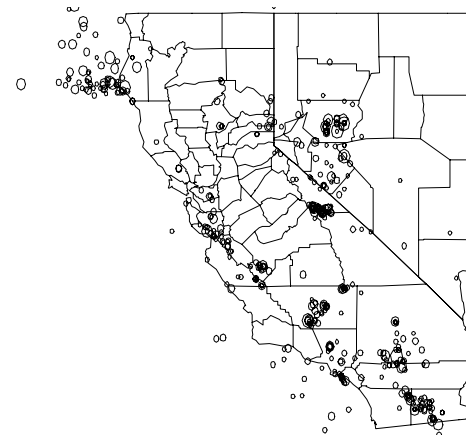


Figure 2. Map of All Longleaf Pines in the 150×120 m Study Region B (Inner Rectangle) and the 30 Meter Wide Guard Region $B_+ - B$. The direction north is toward the right side of the page.

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California Earthquakes



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Point pattern analysis is primarily concerned with modeling the locations of events, for example the locations of:

- Trees
- Birds' nests
- Ants' nests
- Earthquake epicenters
- Cancer cases
- Galaxies

Objectives: Point Pattern Analysis

1. To determine if the point pattern is completely random;
2. If the pattern is not completely random, fit an explanatory point process model to the data.

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Complete Spatial Randomness

Definition: A point pattern is *completely random* if it is realized from a homogeneous Poisson process.

Definition: For a homogeneous Poisson process with intensity λ

1. The number of events (trees) $N(A)$, in a study region A is Poisson distributed with mean $\lambda|A|$

$$\Pr\{N(A) = n\} = \frac{1}{n!} e^{-\lambda|A|} (\lambda|A|)^n$$

2. Conditional on the number of events (trees), the event locations are independently sampled from a uniform distribution on A .

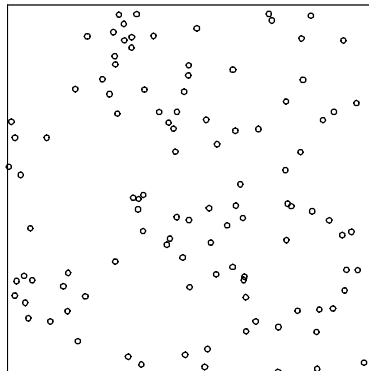
Definition: The *intensity* λ is equal to the mean number of events per unit area.

Note: In ecology, the intensity is called the density. In statistics, we use the term intensity to distinguish it from a probability density function.

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Completely Random Pattern

Complete Spatial Randomness



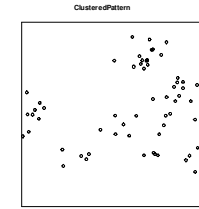
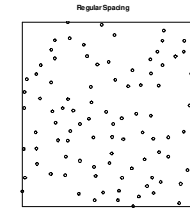
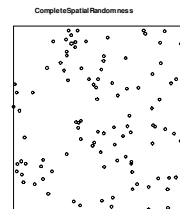
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Complete spatial randomness is the null model against which spatial point patterns are often compared.

Completely Random

Regular

Clustered



In Ecology:

- Regular spacing may result from intraspecific competition for limited resources;
- Clustered patterns may result from:
 - Clustering of offspring around their parents;
 - Response to a heterogeneous environment.

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Ripley's K-Function

Ripley's K-function is the most effective tool for assessing departure from complete spatial randomness.

Definition:

$$K(r) = \frac{\text{Mean number of trees within distance } r \text{ of an arbitrary tree}}{\lambda}$$

Estimation:

$$\hat{K}(r) = \frac{1}{\hat{\lambda}N} \sum_{i \neq j} w_{ij} I(d_{ij} \leq r)$$

where

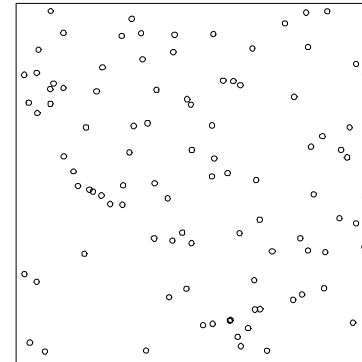
$$\hat{\lambda} = \frac{N}{|A|}$$

is the number of trees in the study region divided by the area of the study region.

What is this?

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Consider the point pattern of trees:

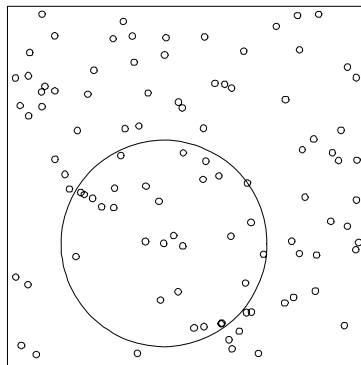


Here there are 100 trees in a 10×10 region. So

$$\hat{\lambda} = \frac{100}{10 \times 10} = 1$$

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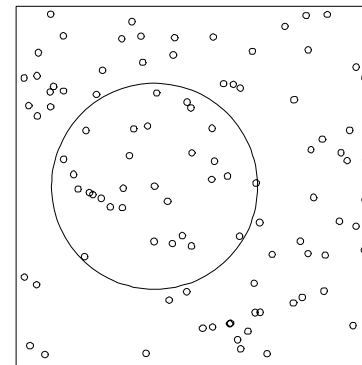
Place a circle of radius r around an arbitrary tree:



Count the number additional of trees within the circle.

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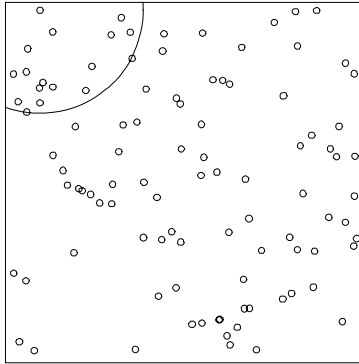
Repeat for each of the remaining trees:



Counting the number of additional trees within each circle.

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Edge Correction



For trees close to the edge of the study region, we cannot observe the number of trees within radius r .

Here, we give the neighboring trees a weight w_{ij} equal to one divided by the portion of the circle of radius d_{ij} inside the study region.

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The results are averaged over all base trees

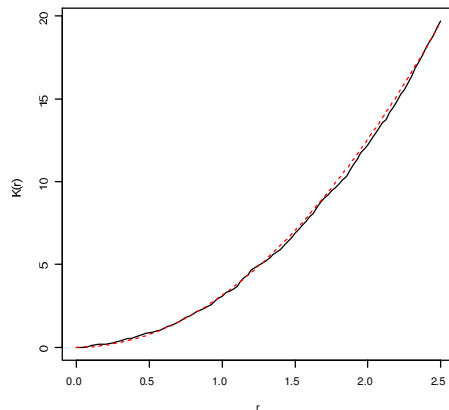
$$\frac{1}{N} \sum_{i \neq j} w_{ij} I(d_{ij} \leq r)$$

and then divided by the estimated intensity $\hat{\lambda}$ to obtain the estimate

$$\hat{K}(r) = \frac{1}{\hat{\lambda} N} \sum_{i \neq j} w_{ij} I(d_{ij} \leq r)$$

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Plot $\hat{K}(r)$ against r



Note: Under complete spatial randomness,

$$K(r) = \pi r^2$$

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Note: Even for strong departures from complete spatial randomness, the difference between the empirical K-function and its expectation under complete spatial randomness is small.

Therefore, a plot of the K-function may not be very informative.

Solution: Linearizing Transformation:

$$L(r) = \sqrt{K(r)/\pi} - r$$

- Under complete spatial randomness

$$L(r) = 0$$

- For clustered patterns

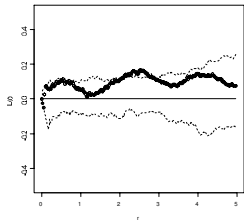
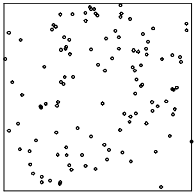
$$L(r) > 0$$

- For regular spacing

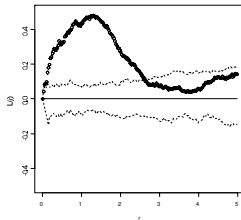
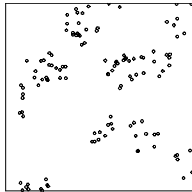
$$L(r) < 0$$

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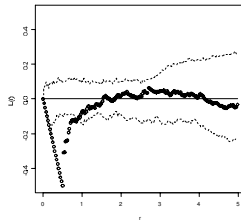
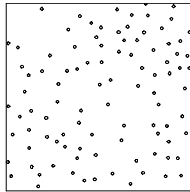
Completely Random



Clustered



Regular



Note: By plotting the L-function against distance, all scales of pattern can be examined.

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Point Process Models Inhomogeneous Poisson Process

Definition: The *intensity* of a point process is

$$\lambda(\mathbf{s}) = \lim_{|ds| \rightarrow 0} \frac{E\{N(ds)\}}{|ds|}$$

The intensity can be viewed as a local density. Regions with high intensities will tend to contain large numbers of trees, while regions with low intensities will tend to contain few trees.

- $N(ds)$ is the number of trees in a small region ds surrounding the location \mathbf{s}
- $E\{N(ds)\}$ is the mean number of trees in ds
- $|ds|$ is the area of the region ds
- Thus, the intensity $\lambda(\mathbf{s})$ is the mean number trees per unit area, as a function of location \mathbf{s} .

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Inhomogeneous Poisson Process

The inhomogeneous Poisson process may be used to model the impact of spatial variation in environmental characteristics (e.g., elevation, light intensity, nutrient concentrations) on a point pattern.

Definition: For an inhomogeneous Poisson process with intensity λ

1. The number of events (trees) $N(A)$, in a study region A is Poisson distributed with mean

$$\Lambda(A) = \int_A \lambda(\mathbf{s}) d\mathbf{s}$$

That is, the probability that the number of events $N(A)$ equal to n is

$$\Pr\{N(A) = n\} = \frac{1}{n!} e^{-\Lambda(A)} (\Lambda(A))^n$$

2. Conditional on the number of events, the event locations are independently sampled from a probability density function proportional to $\lambda(\mathbf{s})$.

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Space-Varying Covariates

Let

$$x_1(\mathbf{s}), x_2(\mathbf{s}), \dots, x_p(\mathbf{s})$$

denote the values of p space-varying covariates at the location \mathbf{s} in the study region A (e.g., elevation, light intensity, nutrient concentrations, etc.).

The impact of these space-varying covariates on a spatial point pattern may be modeled through the intensity function:

$$\lambda(\mathbf{s}; \boldsymbol{\beta}) = \exp\{\beta_0 + \beta_1 x_1(\mathbf{s}) + \beta_2 x_2(\mathbf{s}) + \dots + \beta_p x_p(\mathbf{s})\}.$$

An inhomogeneous Poisson process with the above intensity is called a *modulated Poisson process*.

Reference

Cox, D.R. (1972). The statistical analysis of dependencies in point processes. In P.A.W. Lewis (ed.), *Stochastic Point Processes*, pp. 55-66. New York: Wiley.

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Parameter Estimation

The *maximum likelihood estimator* is obtained by finding $\hat{\beta}$ that maximizes the log likelihood:

$$L(\beta) = \beta' \sum_{i=1}^n \mathbf{x}(s_i) - \int_A \exp\{\beta' \mathbf{x}(s)\}$$

where

- s_1, s_2, \dots, s_n denote the locations of n trees in the study region A .
- $\mathbf{x}(s)$ = vector of covariates at the location s in A .

Problem: This requires that the values of the covariates be observed for:

- All of the trees in the study region.
- All locations in the study region.

The former may be impractical, and the later impossible to obtain.

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Two Approaches:

1. Rathbun (1996) *Biometrics* **52**, 226-242.
2. Rathbun, Shiffman, and Gwaltney (2006) In *Models for Intensive Longitudinal Data*. T.A. Walls and J.L. Schafer (eds.). Oxford.

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Approach 1

- Sample the covariates at a collection of sites

$$\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_m$$

- Use kriging to predict the values of the covariates at the locations of the trees, and at the unsampled sites.
- Substitute predicted values into the log likelihood:

$$\hat{L}(\beta) = n\beta_0 + \beta_1 \sum_{i=1}^n \hat{x}(s_i) - \int_A \exp\left\{\beta_0 + \beta_1 \hat{x}(s) + \underbrace{\frac{1}{2}\beta_1^2(\sigma^2 - \text{var}(\hat{x}(s)))}_{\text{Bias Correction}}\right\} ds$$

- Find $\tilde{\beta}$ that maximizes the approximate log likelihood $\hat{L}(\beta)$.

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Example: Titi Hammock Data Beech-Magnolia Forest in South Georgia

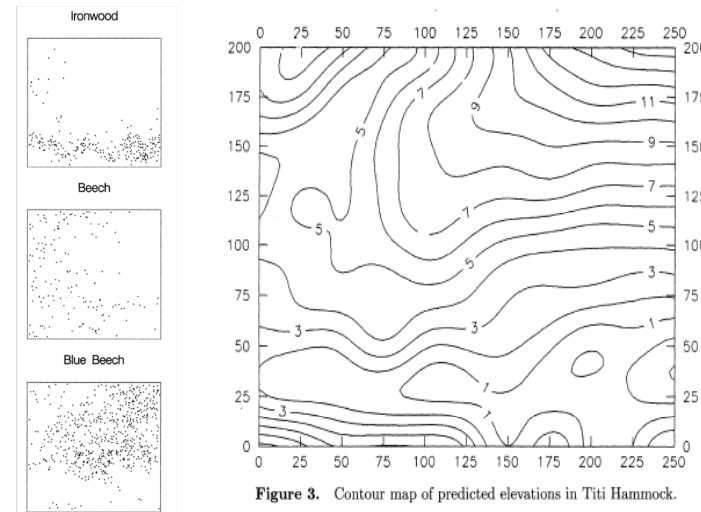


Figure 3. Contour map of predicted elevations in Titi Hammock.

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Results:

Parameter estimates for a modulated Poisson process with intensity (5.2). Standard errors are given in parentheses.

Species	No bias correction		Bias corrected	
	$\hat{\beta}_0$	$\hat{\beta}_1$	$\tilde{\beta}_0$	$\tilde{\beta}_1$
Bay	-4.3729 (0.1355)	-0.9204 (0.0936)	-4.4300 (0.1330)	-0.8819 (0.0883)
Beech	-5.4476 (0.1384)	-0.0613 (0.0262)	-5.4495 (0.1381)	-0.0609 (0.0261)
Blue beech	-5.0711 (0.0821)	0.1362 (0.0113)	-5.0667 (0.0819)	0.1355 (0.0113)
Holly	-5.2169 (0.1099)	0.0161 (0.0183)	-5.2164 (0.1097)	0.0160 (0.0182)
Ironwood	-3.2833 (0.0760)	-0.7621 (0.0432)	-3.3264 (0.0749)	-0.7384 (0.0415)
Magnolia	-5.1454 (0.1135)	-0.0277 (0.0203)	-5.1462 (0.1132)	-0.0276 (0.0202)
Tulip poplar	-4.8605 (0.1750)	-1.0234 (0.1356)	-4.9270 (0.1717)	-0.9725 (0.1265)

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Approach 2

Data Requirements: Covariates are observed

- Locations of the trees

$$\mathbf{s}_1, \mathbf{s}_2, \dots, \mathbf{s}_n$$

- Random locations from the study region

$$\mathbf{u}_1, \mathbf{u}_2, \dots, \mathbf{u}_m$$

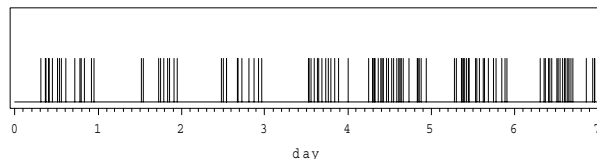
Find $\tilde{\beta}$ that maximizes the approximate log likelihood

$$\hat{L}(\beta) = \beta' \sum_{i=1}^n \mathbf{x}(\mathbf{s}_i) - \frac{|A|}{m} \sum_{j=1}^m \exp\{\beta' \mathbf{x}(\mathbf{u}_j)\}$$

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Example: Ecological Momentary Assessment of Smoking

Times at which cigarettes were lit by a smoker



Time-Varying Covariates

- Negative Affect
- Arousal
- Attention
- Restlessness

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Results

Parameter	Modulate Poisson	
	Estimate	SE
Intercept	-0.05924	0.00839
Negative Affect	0.01950	0.01077
Arousal	-0.01594	0.01078
Attention	-0.01787	0.01198
Restlessness	0.21017	0.01577

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Extensions:

- Obtain covariates on a thinned sample of trees. Visit each tree and sample the covariates with known probability p . More generally, p may depend on location.
- Use alternative designs for covariate sample sites:
 - Stratified Random Sample
 - Transect Samples - Random parallel transects, and random sites along each transect.